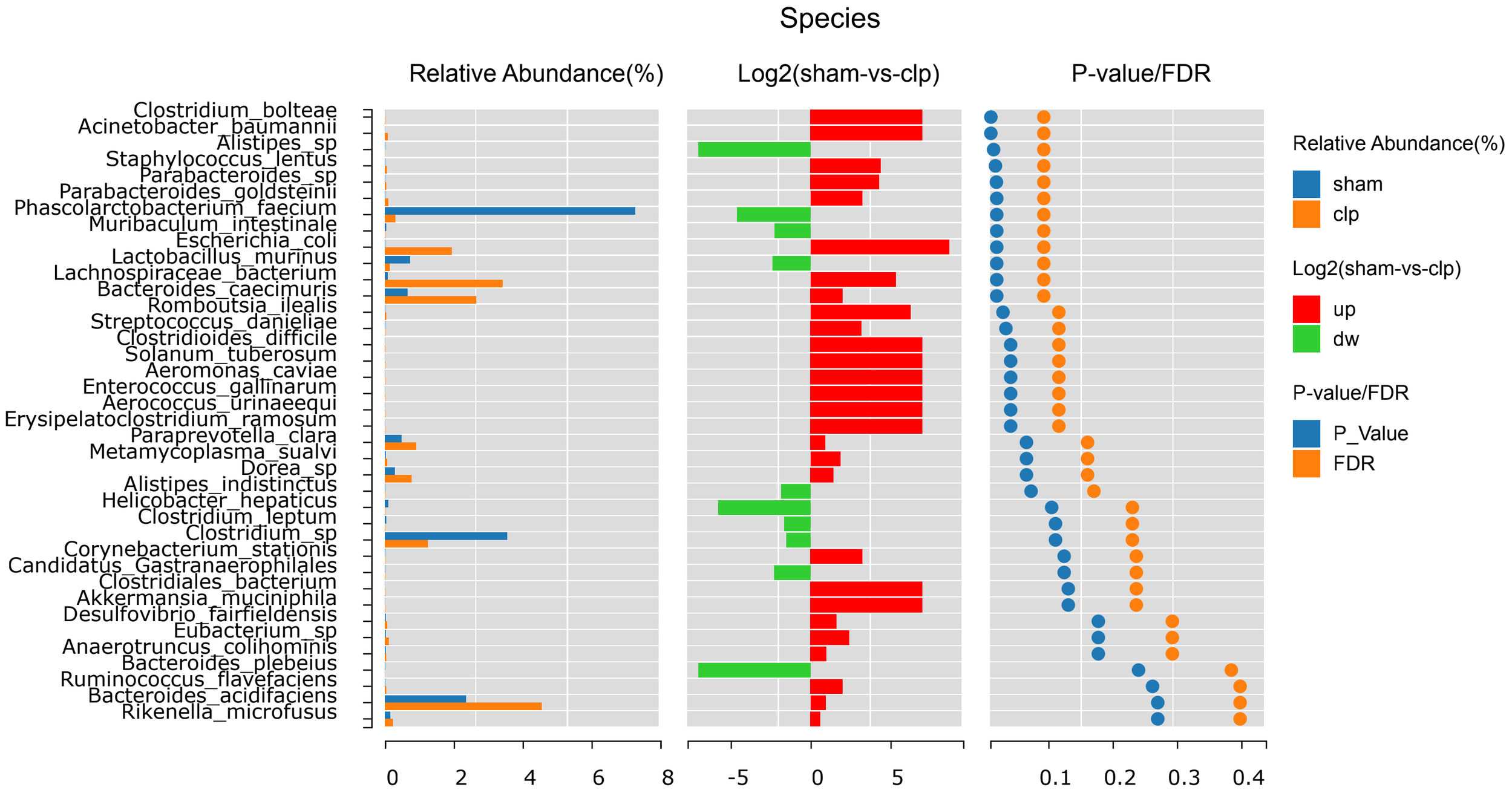
Species
Relative Abundance(%)
Log2(sham-vs-clp)
P-value/FDR
Clostridium_bolteae
Acinetobacter_baumannii
Alistipes_sp
Staphylococcus_lentus
Parabacteroides_sp
Parabacteroides_goldsteinii
Phascolarctobacterium_faecium
Muribaculum_intestinale
Escherichia_coli
Lactobacillus_murinus
Lachnospiraceae_bacterium
Bacteroides_caecimuris
Romboutsia_ilealis
Streptococcus_danieliae
Clostridioides_difficile
Solanum_tuberosum
Aeromonas_caviae
Enterococcus_gallinarum
Aerococcus_urinaeequi
Erysipelatoclostridium_ramosum
Paraprevotella_clara
Metamycoplasma_sualvi
Dorea_sp
Alistipes_indistinctus
Helicobacter_hepaticus
Clostridium_leptum
Clostridium_sp
Corynebacterium_stationis
Candidatus_Gastranaerophilales
Clostridiales_bacterium
Akkermansia_muciniphila
Desulfovibrio_fairfieldensis
Eubacterium_sp
Anaerotruncus_colihominis
Bacteroides_plebeius
Ruminococcus_flavefaciens
Bacteroides_acidifaciens
Rikenella_microfusus
0
2
4
6
8
-5
0
5
0.1
0.2
0.3
0.4
Relative Abundance(%)
sham
clp
Log2(sham-vs-clp)
up
dw
P-value/FDR
P_Value
FDR